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Minimum
Maximum
                                                                                                                                                                                                 Database
                                                                                                                                                                                                                      Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                             Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                   Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                 Searched:
                                                                                                                                                                                                                                                                                                                                                                                       Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OM protein -
SPTREMBL_15:*

1: sp_archea:*
2: sp_bacterie
3: sp_fungi:*
4: sp_human:
5: sp_human:
6: sp_mamme
7: sp_mhc.
8: sp_org
9: sp_ph
10: sp_
11: sp
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14:
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                                                                                                                                                                                                                                                                             seq
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length: 2000000000
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Gapop 10.0 ,
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384
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                                                                                                                                                                                                                                                                                                                                                                                                                  NRQRDPQQQYEQCQERCQRH......EEQQREDEEKYEERMKEEDN
                                                                                                                                                                   sp_archea:*
sp_bacteria:*
                                                                                    sp_organelle:*
                                                                                                                          sp_invertebrate:*
     sp_vertebrate:*
sp_unclassified:*
                             sp_rodent:*
sp_virus:*
                                                         sp_plant:*
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26.982 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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19	18	17	16	15	14	13	12	11	10	9	8	7	6	5	4	w	2		Result
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26.2	26.4	26.7	27.0	27.1	27.2	27.3	27.3	28.5	28.9	30.1	30.5	30.5	31.1	43.0	45.6	93.5	95.6	100.0	Query Match I
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25.3	25.3	25.3	25.3	25.5	25.5	25.5	25.7	25.7	25.7	5	25.7	25.7	25.7	25.7	5	5	5	5	5	<u>ب</u>	ū	5	5	25.9	5
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Q9und7 homo sapien			015410 homo sapien	035788 rattus norv	Q9vsk3 drosophila	dicty	homo	homo	homo	homo	09uke0 homo sapien	Q9uke3 homo sapien	homo	Q9uke1 homo sapien	Q9ukd8 homo sapien			095033 homo sapien	0 mus	075172 homo sapien	095819 homo sapien	Q9nst7 homo sapien	Q22540 caenorhabdi	Q9u9s7 dictyosteli	Q9vva9 drosophila

ALIGNMENTS

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RESULT
OPERATE
RESULT
Q9SPL5
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Best Local Similarity
Matches 69; Conserv
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Q9SPL4;
Q1-MAY-2000 (TremBLrel. 13,
Q1-MAY-2000 (TremBLrel. 13,
Q1-OCT-2000 (TremBLrel. 15,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE-NUT KERNEL;
Marcus J.P., Goulter K.C., Green J.L., Manners J.M.;
Marcus J.P., Goulter K.C., Green J.L., Manners J.M.;
"A family of antimicrobial peptides is produced by puglobulin protein in Macadamia integrifolia.";
Plant J. 0:0-0(1999).
EMBL; AF161884; AAD54245.1; -.
HSSP; P02853; 2PHL.
HTERPRO; IPRO01113; -.
PTAM; PF00546; Seedstore 7s; 1.
PFAM; PF00546; Seedstore 7s; 1.
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Eukaryota; Viridiplantae; Embryophyta; Tracheophyta;
Magnoliophyta; eudicotyledons; Proteaceae; Macadamia.
NCBI_TaxID=60698;
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                                                                                                                                                                                                                                                                                   61 EERMKEEDN 69
                                                                                                                                             EERMKEEDN 185
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Last annotation update)
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Pred. No. 5.4e-27;
); Mismatches 0;
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RESULT
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Best Local S
Matches 65
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Best Local S
Matches 66
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01-MAY-2000
01-MAY-2000
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01-MAY-2000
01-OCT-2000
                                                                                                                                                                                                                                                                                                                                                             TISSUE=NUT KERNEL;
Marcus J.P., Goulter K.C.,
"A family of antimicrobial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q9SPL5;
                                                                                                                                                                                           PFAM; PF00546; Seedstore_7s;
NON_TER 1 1
SEQUENCE 625 AA; 73586 MW
                                                                                                                                                                                                                                                                              HSSP; P02853;
                                                                                                                                                                                                                                                                                                    EMBL; AF161885; AAD54246.1;
                                                                                                                                                                                                                                                                                                                    globulin protein in Macadamia
Plant J. 0:0-0(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Magnoliophyta; eu
NCBI_TaxID=60698;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Macadamia integrifolia (Macadamia nut).
Eukaryota; Viridiplantae; Embryophyta; Tracheophyta;
Magnoliophyta; eudicotyledons; Proteaceae; Macadamia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VICILIN PRECURSOR
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PFAM; PF00546; Seedstor
SEQUENCE 666 AA; 782
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Macadamia integrifolia (Macadamia nut).
Eukaryota; Viridiplantae; Embryophyta; Tra
Magnoliophyta; eudicotyledons; Proteaceae;
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    76
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NRQRDPQQQYEQCQERCQRHETEPRHMQTCQQRCERRYEKEKRKQQKRYEEQQREDEEKY 60
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                                                                                     65; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                          853; 2PHL.
IPR001113;
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                                                                                     Conservative
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AA; 78217 MW;
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                                                                                                                                                                                             73586 MW;
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94.2%;
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95.7%;
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13,
                                                                                                                                                                                                                                                                                                                                          Green J.L., Manners J.M.; peptides is produced by p mia integrifolia.";
                                                                                   Pred. No. 8.5e
1; Mismatches
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Pred. No. 1.7e-25;
1; Mismatches 2
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8.5e-25;
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eae; Macadamia
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Matches 33
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043358; 01-NOV-1996 (TrEMBLrel. 0
01-NOV-1996 (TrEMBLrel. 0
01-OCT-2000 (TrEMBLrel. 1
                                                                                                                                                                                                                                                                                                                                                                                                Q9SEW4 PRELIMINARY; PRT; 593 AP
Q9SEW4;
Q1-MAY-2000 (TrEMBLrel. 13, Created)
Q1-MAY-2000 (TrEMBLrel. 13, Last sequence u
Q1-OCT-2000 (TrEMBLrel. 15, Last annotation
VICILIN-LIKE PROTEIN PRECURSOR (FRAGMENT).
                                                                                                                                                                                                                                                            Juglans regia (English walnut).

Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids I;
Fagales; Juglandaceae; Juglans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Signal.
                                     SEQUENCE FROM N.A.
STRAIN-CV. SUNLAND; TISSUE-SOMATIC EMBRYO LINE;
Teuber S.S., Jarvis K.C., Peterson W.R., Dandekar A.M., Ansari A.A.;
"Identification and cloning of a cDNA encoding a vicilin-like protei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              McHenry L., Fritz P.J.;
"Comparison of the structure and nucleotide sequences of vicilin of cocoa and cotton raise questions about vicilin evolution.";
plant Mol. Biol. 18:1173-1176(1992).
                                                                                                                                                                                                                      Fagales; Juglandaceae;
NCBI_TaxID=51240;
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EMBL; X62625; CAA44493.1; -.
EMBL; X62626; CAA44494.1; -.
HSSP; P02853; 2PHL.
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Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta; Eukaryota; Viridiplantae; Embryophyta; Eudoicotyledons; core eudicots; Rosidae; eurosids II; Majvales; Malvaceae; Theobroma.
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Pred. No. 1.7e-08;
9; Mismatches 14
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01-MAY-2000
 Q9ZWI3;
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                                                                                                                                                             Plasmid
                                                                                                                                                                   Rieben W.K., Gonzales C., Go
Hughes J.E., Welker D.L.;
Submitted (APR-1997) to the
EMBL; AF000580; AAC14390.1;
                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                        MEDLINE-9819836; PubMed=9539429;
Rieben W.K. Jr., Gonzales C.M., Gonzales S.T.,
Kiyosawa H., Hughes J.E., Welker D.L.;
"Dictyostelium discoideum nuclear plasmid Ddp5
the Ddp1 and Ddp2 plasmid families.";
Genetics 148:1117-1125(1998).
                                                                                                                                                                                                                                                                                                              Dictyostelium Plasmid Ddp5.
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Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF066055; AAF18269.1; ..
HSSP; P02853; 2PHL.
INTERPRO; IPR0011113; ..
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                                                                                                                                                                                                                                                                                                       Eukaryota; Dictyosteliida; Dictyostelium
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38; Conservative
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         PRELIMINARY;
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Pred. No. 1.4e
9; Mismatches
                                                                                                             Score 119.5; DB 5;
Pred. No. 0.0025;
2; Mismatches 18;
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01-JAN-1998
01-JUN-2000
                                                                                                                                                                                                         "Cell-homologous genes in the Kaposi's sarcoma-associated rhadinovirus human herpesvirus 8: determinants of its pathogenicity?";

J. Virol. 71:4187-4192(1997).

EMBL; U93872; AAB62657.1; ...
INTERPRO; IPR002017. ...
                                                                                                                                                                                            INTERPRO;
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               040947;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cucurbita maxima (Pumpkin) (Winter squash).
Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
Magnollophyta; endicotyledons; core eudicots; Rosidae; eurosids I;
Cucurbitales; Cucurbitaceae; Cucurbita.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Multiple functional proteins are produced by cleaving of a single precursor by vacuolar processing enzyme."; J. Biol. Chem. 274:2563-2570(1999).
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                                                                                                                                                                                                                                                                                                                                                                                        Viruses; dsDNA viruses, no RNA stage; Gammaherpesvirinae; Rhadinovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                       Kaposi's sarcoma-associated herpesvirus (KSHV) (Human herpesvirus
                                                                                                                                                                                                                                                                                                                                                                                                                                               ORF 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INTERPRO; IPR0011113; -.
PFAM; PF00546; Seedstore_7s;
PRODOM; PD081059; -; 1.
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                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=37296;
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                                    589
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 62
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                                                     RQRDPQQQYEQCQERCQRHETEPRHMQTCQQRCERRYEKEKRKQQKRYEEQQREDEEKYE
ERMKEED
                               QQQDEQQQDEQQQDEQEQQDEQQQDEQQQDEQEQQEEQEQQEEQEQQEEQEQQEEQE 648
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                                                                                                        Similarity
21; Conserv
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25; Conser
                                                                                                                                                                                              1089 AA;
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                                                                                                        Conservative
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 68
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Pred. No. 0.0043;
6; Mismatches 20;
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Pred. No. 0.0034;
4; Mismatches 29
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                                                                                                                                                                                          48AE2F9837E1AB6A CRC64;
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RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Berson K.Y., Benos P.V., Berman B.P., Brokstein P., Brottier P.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier A., Chandra I.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W.,
RA Dodson K., Doup L.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Posler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Alodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D., Kraft C., Kravitz S., Kulp D., Lai Z.,
Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
Liasko P., Lei Y., Levitsky A.A., Li J., McPherson D.,
Lako P., McPherson D.,
Lako P., McPherson D., McPherson D., Melson D.R., Mels
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Best Local
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Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,
Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
Shue B.C., Siden Klamos I., Simpson M., Skupski M.P., Smith T.,
Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
Sylirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,
Wang Z. Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
Wang Z. Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
Williams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A.,
Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L.,
Zheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu X., Smith H.O
Glibbs R.A., Myers E.W., Rubin G.M., Venter J.C.,
"The Action of the Parket Processing P
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01-MAY-2000
01-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "The genome sequence of Drosophila melanogaster."; Science 287:2185-2195(2000).
EMBL: AE003588; AAF51469.1; -
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SEQUENCE
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Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muso
Ephydroidea; Drosophilidae; Drosophila.
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MEDLINE=20196006;
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ORDPOQQYEQCQERCQRH----ETEPRHMQTCQQRCERRYEKEKRKQQKRYEEQQREDEE 58
                                                                                                                                                         26;
                                                                                                                                               Similarity 36.0
26; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IPR001304; -.
                                                                                                                                                                                                                                                                                                                                                                                                        554 AA;
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Last annotation update)
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                                                                                                                                                                                                 Score 115.5; DB Pred. No. 0.0034;
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Best Local
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Parry J.P., Peruzzi D., Edelin
Submitted (MAY-1997) to the I
EMBL; U52064; AAC55944.1; -.
EMBL; U75698; AAC57158.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=97048116; PubMed=8892957; Cesarman E., Nador R.G., Bai F., Bo Moore P.S., Chang Y., Knowles D.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kaposi's sarcoma-associated herpesvirus (KSHY) (Human herpesvirus Viruses; dsDNA viruses, no RNA stage; Herpesviridae; Gammaherpesvirinae; Rhadinovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         receptor and cyclin D homologs which and malignant lymphoma.";
J. virol. 70:8218-8223(1996).
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01-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Q98148;
01-FEB-1997
                                                                                                                                                                                                                                                                  Submitted
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Parry J.P.,
                                                                                                                                                                                                                                                                                                                                                              Russo J.J.,
Parry J.P.,
                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A. MEDLINE=97121480; PubMed=8962146;
                                                                                                                                                                                                                                                                                                                                                                                                                                     genes by
                                                                                                                                                                                                                                                                                                                                                                                                                                             Moore P.S., Bashoff C., Weiss R.A., Chang Y.; "Molecular mimicry of human cytokine and cytokine response
                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=97094384; PubMed=8939871;
Moore P.S., Bashoff C., Weiss R.A
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                                                                                                                                                                                                                                                                                                                                                   "Nucleotide sequence of the Kaposi sarcoma-associated
                                                                                                                                                                                                                                                                                                                                                                                                                           Science
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                                                                                              . Similarity
21; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                         y KSHV.";
274:1739-1744(1996).
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                                                                                                                                                                                                                                                                (OCT-1996)
                                                                                                                                                                     IPR002017;
                                                                                                                                                         1162
                                                                                                                                                                                                                                                                           Bohenzky R.A.,
Peruzzi D., Ed
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                                                                                                                                                                                                                                                                                                                            Acad.
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Peruzzi D., Edelman I.S.,
                                                                                              Conservative
                       68
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                                                                                                                                                                                                                   Edelman
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                                                                                                                                                                                                                                                                           Chien M.C.,
elman I.S.,
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Last sequence update)
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                                                                                                       Score 111; DB 1
Pred. No. 0.016;
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                                                                                              Mismatches
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Chang Y., Moore P.S.;
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Chang Y., Moore P
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).S.;
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Matches 22; Conserv
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INTERPRO; IPR000533; -.
INTERPRO; IPR0002017; -.
INTERPRO; IPR002965; -.
INTERPRO; IPR002965; -.
PRINTS; PR010194; TROPOMYOSIN.
PRINTS; PR01217; PRICHEXTENSN.
SEQUENCE 1129 AA; 131346 MW;
                                                                                                                                                                                                                                                        01-NOV-1998
01-NOV-1998
01-JUN-2000
                                                                                    PFAM; PFC
SEQUENCE
                                                                                                                 "Dictyostelium TRFA homologous growth and early development.";
J. Biol. Chem. 0:0-0(1998).
EMBL; AB009080; BAA33143.1;
                                                                                                                                                                                                                                                                                         077033;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kaposi's sarcoma-associated Viruses; dsDNA viruses, no I Gammaherpesvirinae; Rhadinov
                                                                                                                                                             Saito J., Adachi H.,
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                                                                                                                                                                                                     NCBI_TaxID=44689;
                                                                                                                                                                                                                 Eukaryota;
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                                                                                                          INTERPRO;
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RPRO; IPR001440;
; PF00515; TPR; 9
ENCE 1390 AA;
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                                 16;
                                           Similarity
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                                Conservative
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160421 MW;
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                                          27.3%;
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14,
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Pred. No. 0.02:
25; Mismatches
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Pred.
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                                Mismatches
                                                                                   BAAB9926656002DE CRC64;
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No. 0.
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Best Local
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                              P91152
P91152;
01-MAY-1997
01-MAY-1997
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SIMILAR TO D
C43E11.3
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Q9TW28;
01-MAY-2000
01-MAY-2000
01-OCT-2000
        Eukaryota; Metazoa;
                     Caenorhabditis elegans
                                                                                                                                                                                                                                                                                        PROSITE; PS00152; ATPASE_ALPHA_BETA; UNKNOWN_1.
PROSITE; PS50003; PH_DOMAIN; 1.
SEQUENCE 1737 AA; 195860 MW; 39CBF9C08EFEF8
                                                                                                                                                                                                                                                                                                                                                                                                         Dictyostelium discoideum.";
Submitted (SEP-1998) to the EMBL/GenBank/DDBJ databases
EMBL; AB017910; BAA84604.1;
EMBL; AF090533; AA047903.1; -.
HSSP; P08799; 1MND.
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"Dictyostelium discoideum gene for MyoM.";
Submitted (SEP-1998) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                 PFAM; PF00169; PH; 1. PFAM; PF00621; RhoGEF; 1
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Eukaryota; Dictyosteliida;
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                                                                                                                                                                                                                                                                                                                                                     PFAM; PF00063; myosin_head;
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                                         7 (TrEMBLrel. 03, Created)
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DROSOPHILA SU(VAR)3-9 PROTEIN.
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                                                                                              PRELIMINARY;
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 Peloderinae;
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           Nematoda;
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39.1%;
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a; Dictyostelium
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oda; Chromadorea;
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Pred. No. 0.
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           Rhabditida; Rhabditoidea;
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O9YOC9;
O1-NOV-1999 (TrEMBLrel. 12, Created)
O1-NOV-1999 (TrEMBLrel. 12, Last sequence update)
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Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.
EMBL; U80437; AAB37626.1; ..
INTERPRO; IPR001202; ..
INTERPRO; IPR001214; ..
PFAM; PF00397; WW; 1.
PFAM; PF00396; SET; 1.
PFAM; PF000856; SET; 1.
PROSITE; PS01159; WW_DOMAIN_1; 1.
PROSITE; PS01159; WW_DOMAIN_2; 1.
SEQUENCE 1590 AA; 181719 MW; 084E00974199EB1C CRC64;
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Bonfield J., Burton J., Connell M., Copsey T., Cooper J., Coulson A.,
Craxton M., Dear S., Du Z., Durbin R., Favello A., Fulton L.,
Gardner A., Green P., Hawkins T., Hillier L., Jier M., Johnston L.,
Jones M., Kershaw J., Kirsten J., Laister N., Latreille P.,
Lightning J., Lloyd C., Mcmurray A., Mortimore B., O'Callaghan M.,
Parsons J., Percy C., Rifken L., Roopra A., Saunders D., Shownkeen R.,
Smaldon N., Smith A., Sonnhammer E., Staden R., Sulston J.,
Thierry Mieg J., Thomas K., Vaudin M., Vaughan K., Waterston R.,
Watson A., Weinstock L., Wilkinson Sproat J., Wohldman P.;
"2.2 Mb of contiguous nucleotide sequence from chromosome III of C.
elegans.";
Lee S., Parent C.A., Insall R., Firtel R.A.;
"The novel ras interacting protein required for chemotas signal relay in Dictyostelium.";
Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases EMBL; AF159241; AAD43567.1;
SEQUENCE 838 AA; 94059 MW; EAAD911294226808 CRC64;
                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                            Dictyostelium discoideum (Slime mold).
Eukaryota; Dictyosteliida: Dictyosteliida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gattung S., Scheet P.;
Submitted (JAN-1997) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                          Eukaryota; Dictyostellida; Dictyostellum.
NCBI_TaxID=44689;
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[2]
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36.1%;
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 EERMKEE 67
                    147
                                                              Conservative
                                                                      27.1%; 23.9%;
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